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Fig.1A.

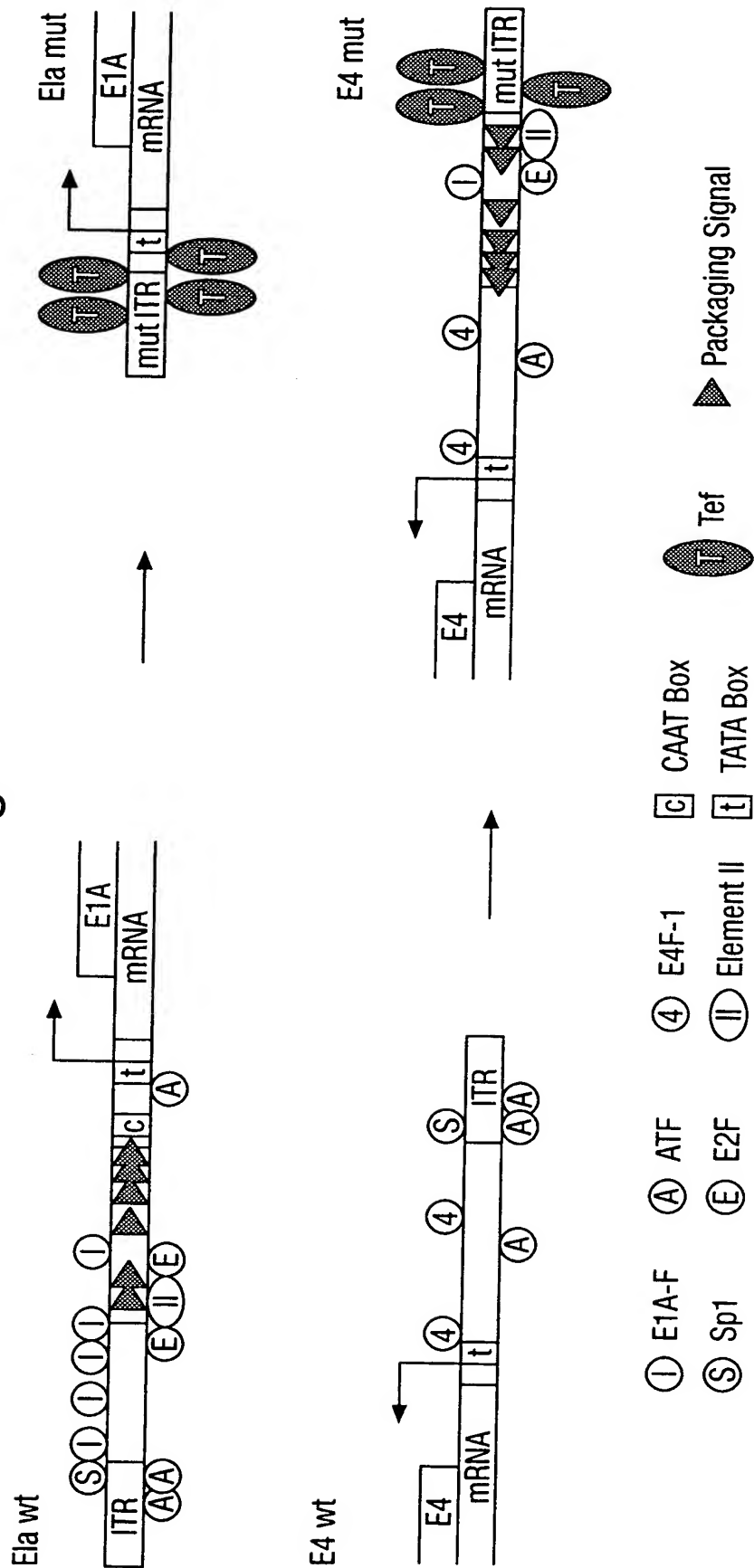
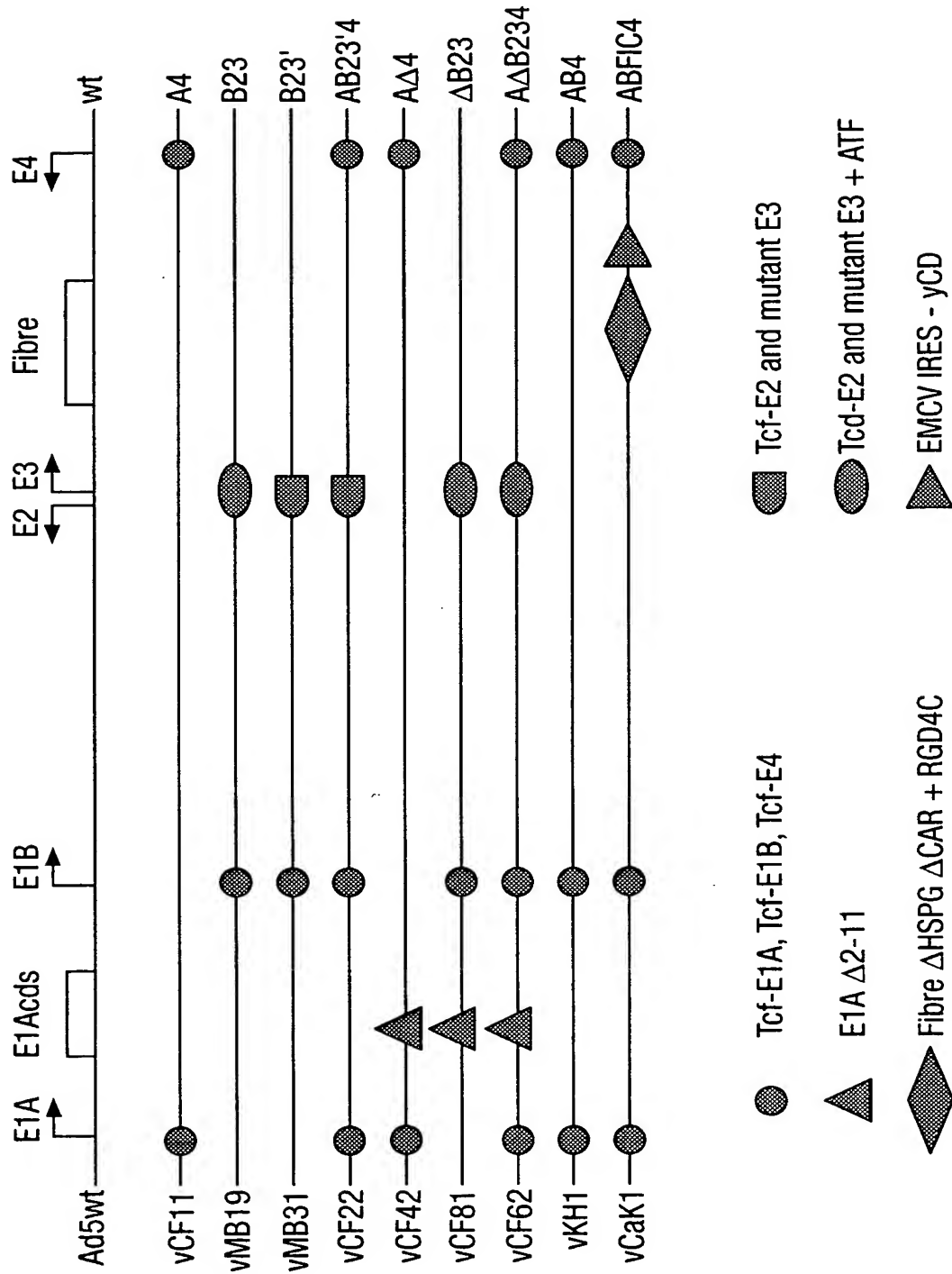




Fig.1B.



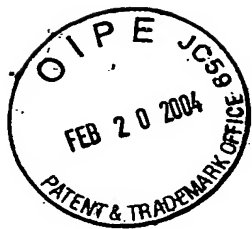


Fig.2.

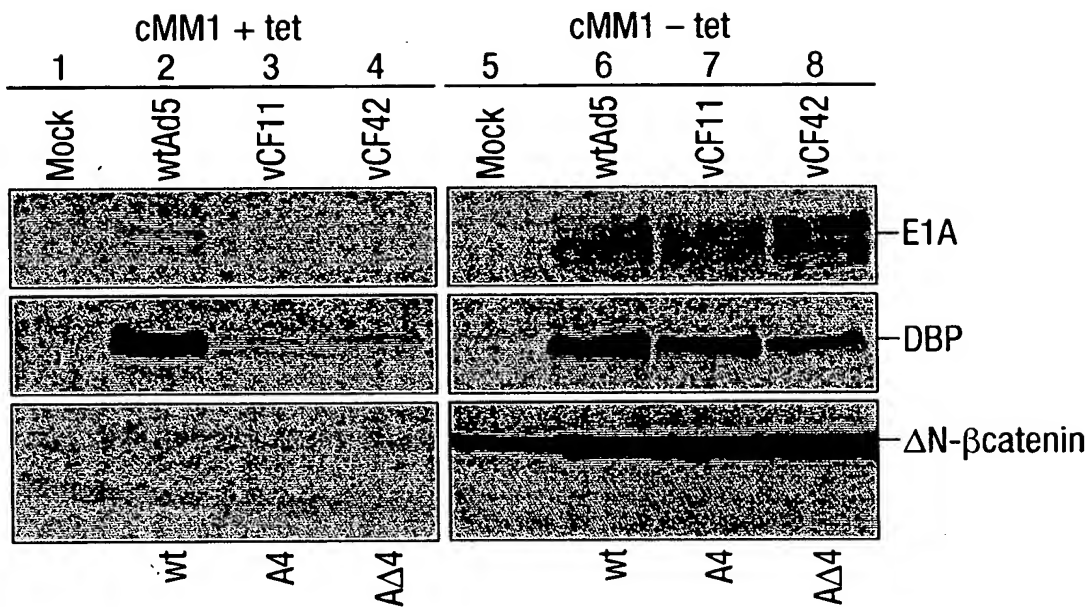
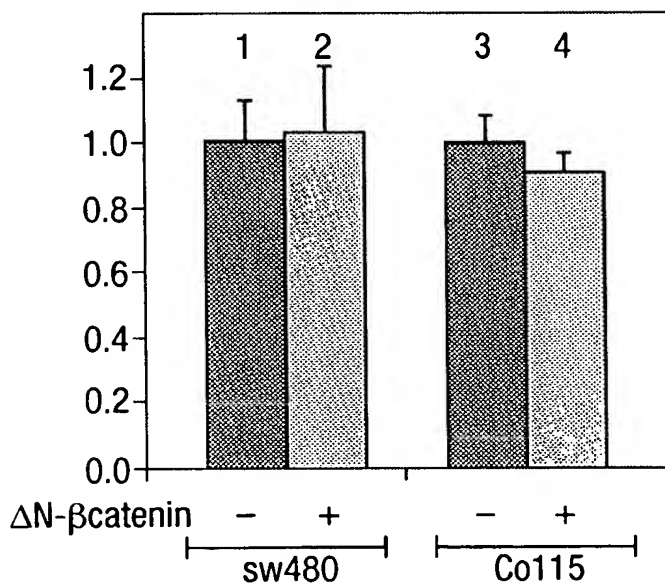
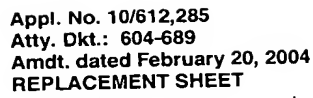


Fig.4.





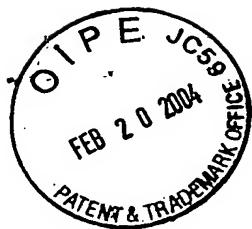


Fig.5A.

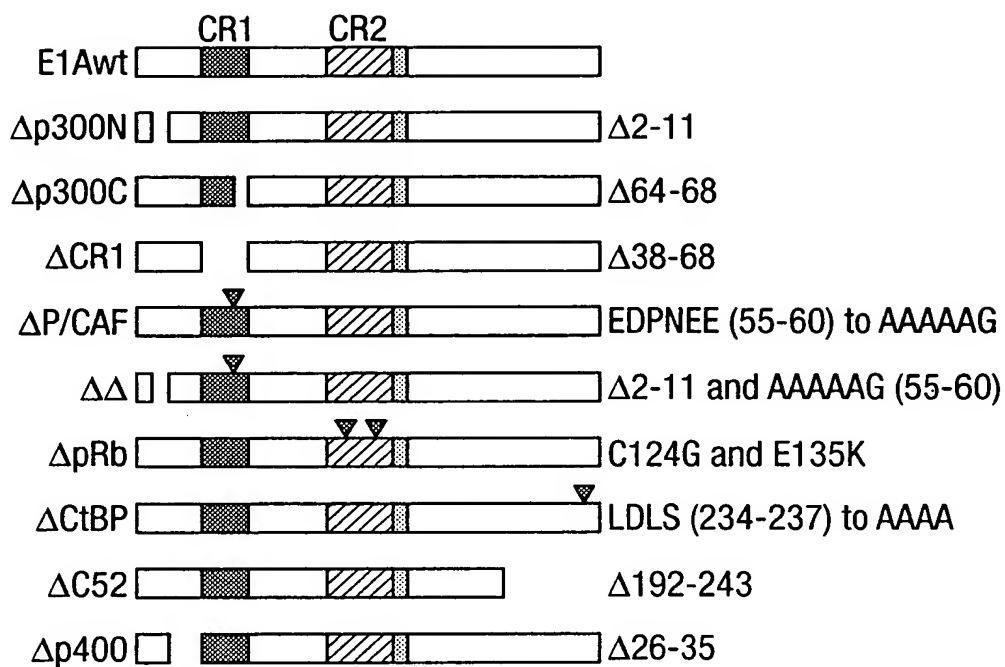


Fig.5B.

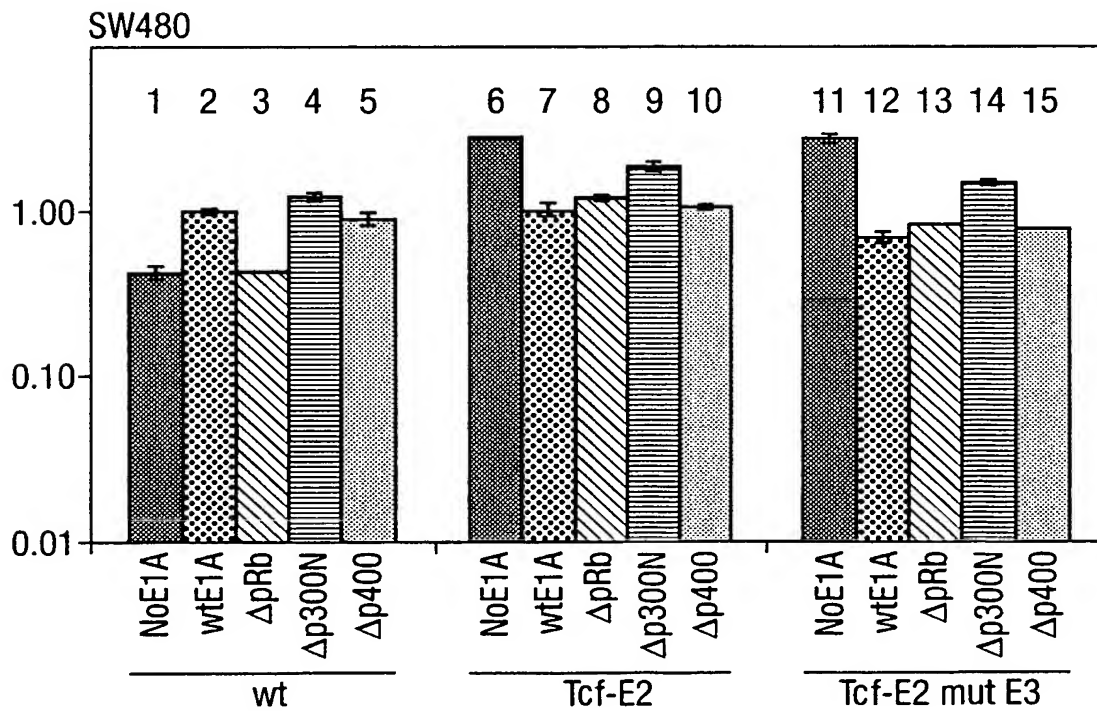




Fig.5C.

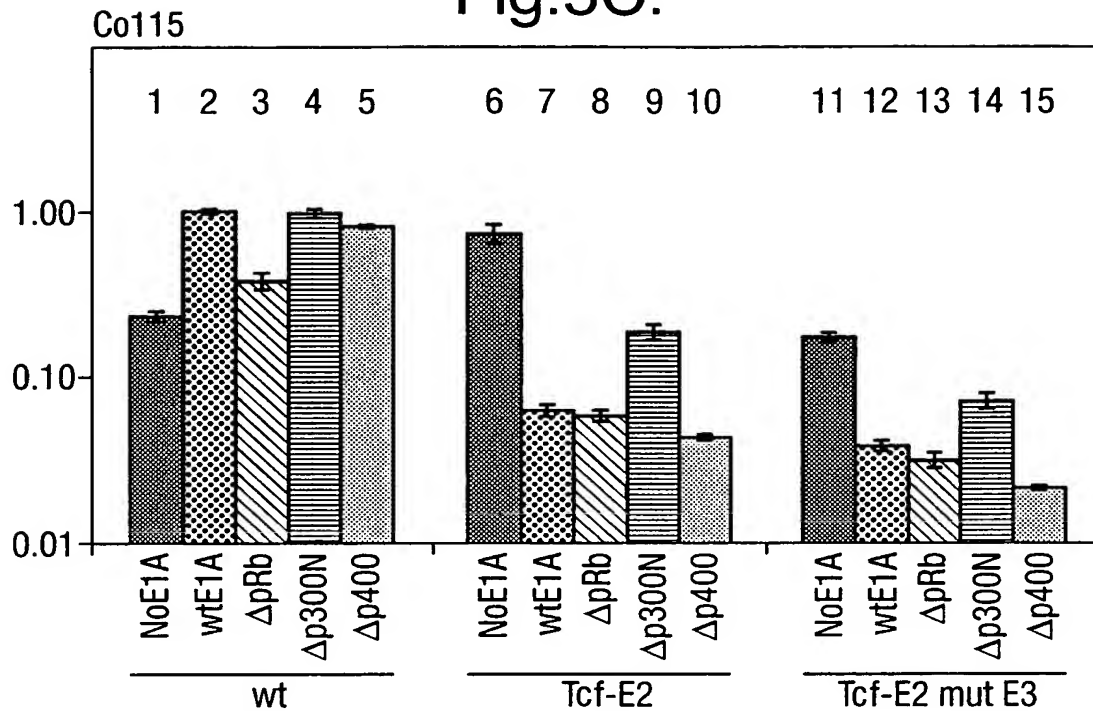
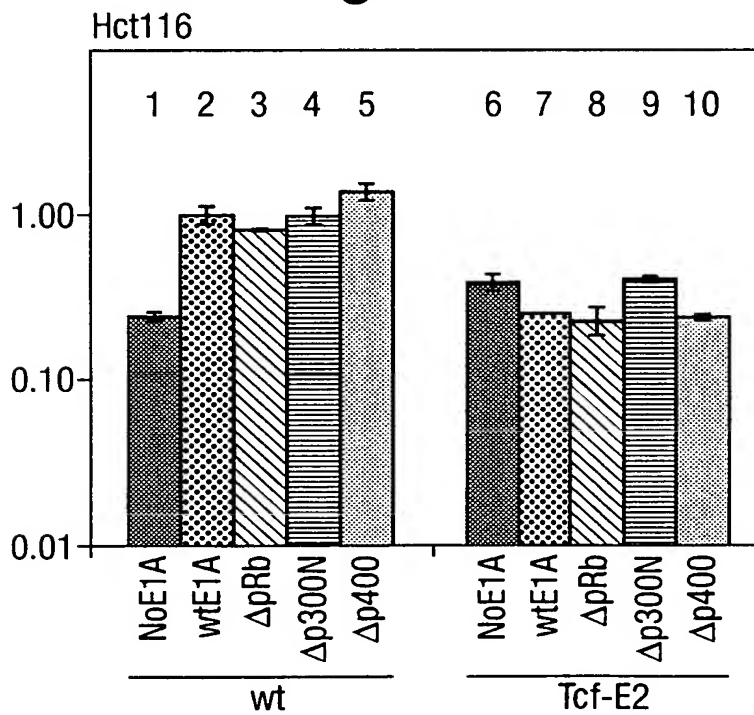


Fig.5D.



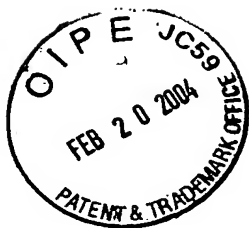


Fig.6A.

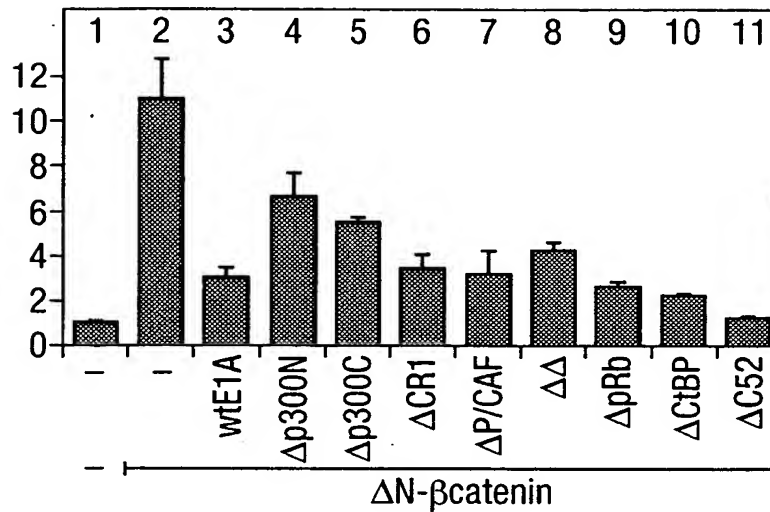


Fig.6B.

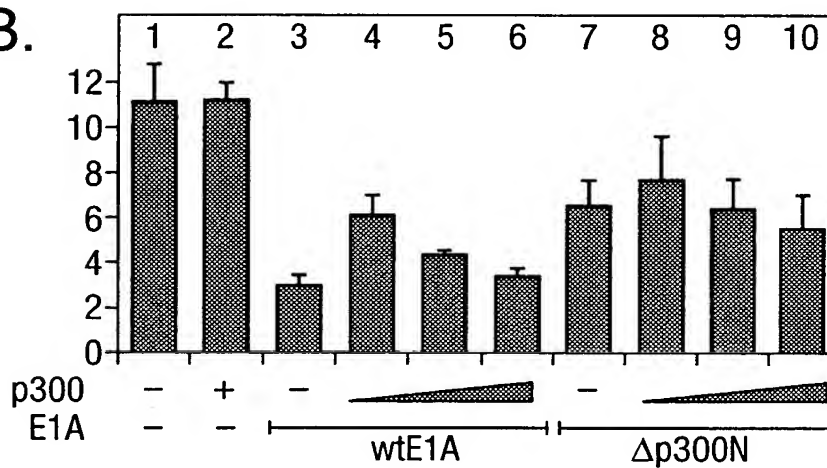


Fig.6C.

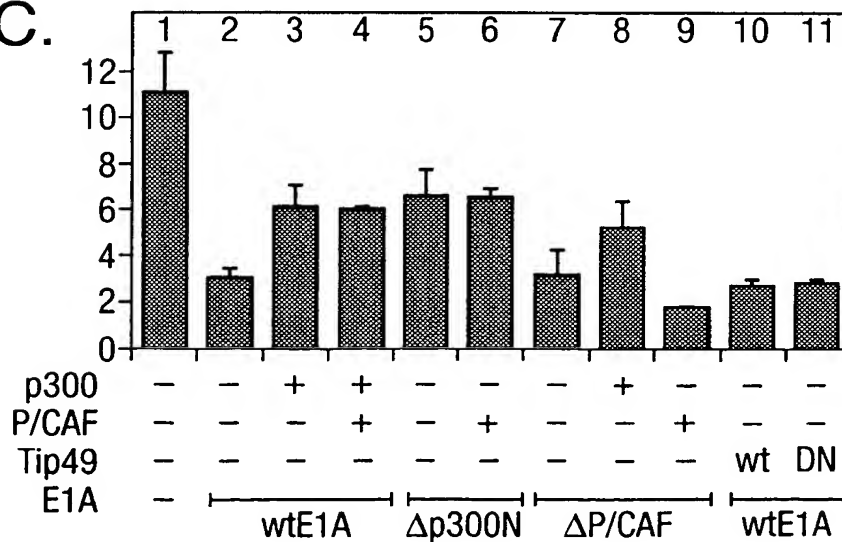






Fig.7A.

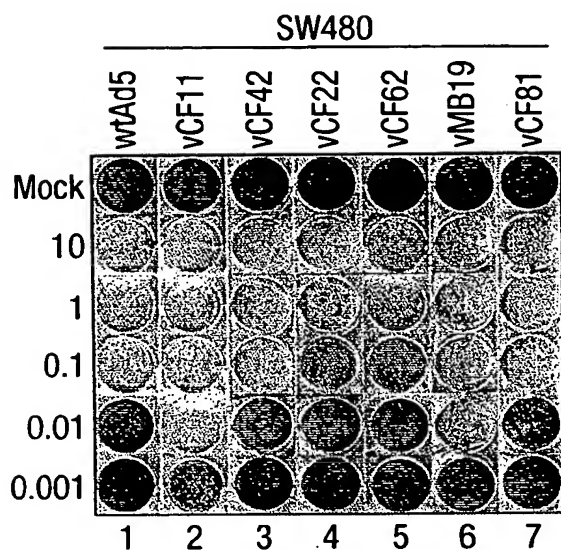


Fig.7B.

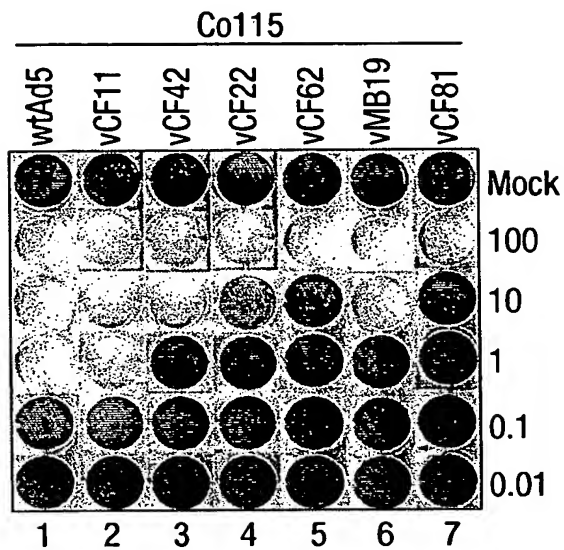


Fig.7C.

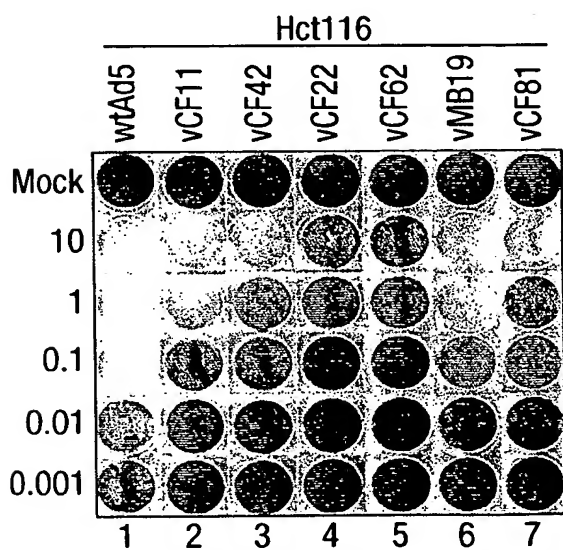


Fig.7D.

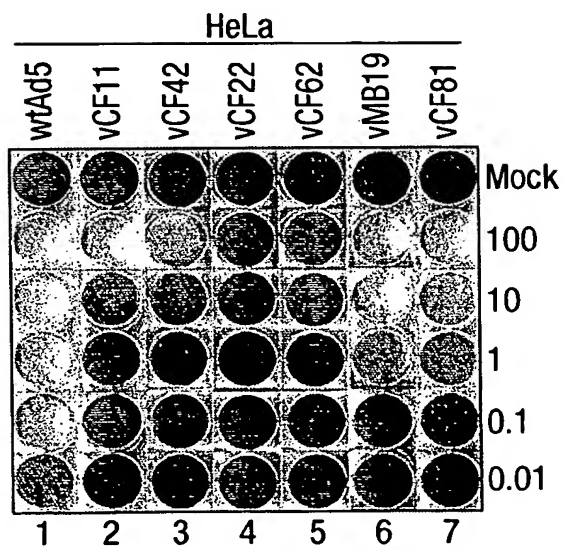
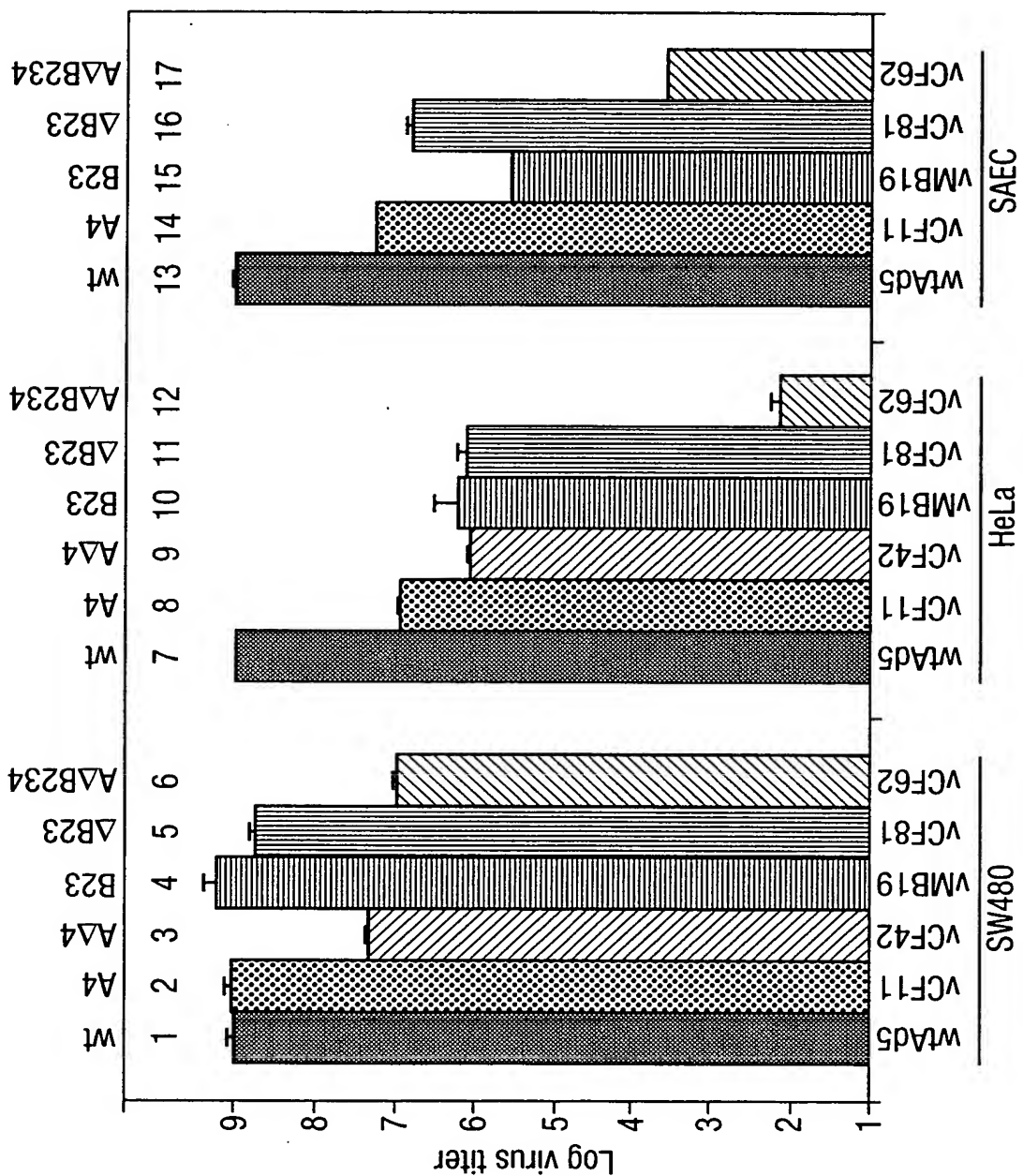
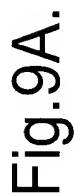




Fig.8.





### Region of max DNA replication

**Sp1** **ATF**

CATCATCAATAATATACCTTATTTTGGATTGAAGCCCAATATGATATGAGGGGGTGGAGTTTGTGACGTGGCGCGGGCGGTG

**CATCATCAATAATACCTTATTTTGGATTGAAGCCAATATGATATGAGGGGGTGGCCCTTTTGATCTTAAATCCCTTTGATC**

Tcf Tcf

ITR - >|

**Spl ATF**

**E1A-F**

**E1A-F**

WT GGAAACGGGGGGTACGTAGTGTGGCGGAAGTGTGATGTTGCAAGTGTGGCGGAACACATGTAAAGCGGATGTGGC

Mut TGGATCCCTTTGATCTCCAACCCTTGTGATCTAGTC

**Tcf**

New ITR ->

**E1A-F**

## E2F

**E1A-F**

Wt AAAAGTGACGTTTTTGGTGTGCGCGCGGTGTACACAGGAAGTGACAAATTTTCGCGCGGTTTAGCGCGATGTTGTAGTAAATT

Mut ::



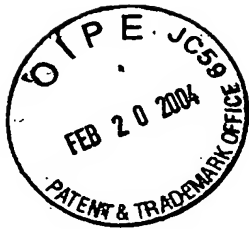


Fig.10A.

E4 promoter

	Cap site	Tata Box	E4F1
Wt	GTCA	<u>CAGTAAAGGGCC</u> AAGTGCAGAGCGAGTATATATAGGACTAAATAATGACGTAA	<u>CGGTTAAAGTCCACAAAAA</u>
Mut	GTCA	<u>CAGTAAAGGGCC</u> AAGTGCAGAGCGAGTATATATAGGACTAAATAATGACGTAA	<u>CGGTTAAAGTCCACAAAAA</u>

ATF

Wt	CACCCAGAAAACCGCACGCGAACCTACGCCCAGAAACGAAAGCCAAAAAACCACAACTTCCTCAAAATCGTCACTTCCGTTT
Mut	CACCCAGAAAACCGCACGCGAACCTACGCCCAGAAACGAAAGCCAAAAAACCACAACTTCCTCAAAATCGTCACTTCCGTTT

E4F1

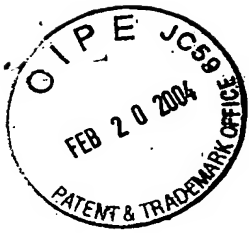
Wt	TCCCACGTTACGTCACTTCCCATTTTAAAGAAAACACTACAATTCCCAACACATACAAGTTACTCCGCCCTAAAC
Mut	TCCCACGTTACGTCACTTCCCATTTTAAAGAAAACACTACAATTCACTAGCAAAACACCTGGCGAGTCTCCACGTAAACGGT

AVII AVI

Wt	CAAAGTCCCCGGCCCTAGACAAAATATTACGGCGTATGAGTAACACAAAATTATTTCAGATTTCACCTTCCCTTATTTCAGTT
Mut	CAAAGTCCCCGGCCCTAGACAAAATATTACGGCGTATGAGTAACACAAAATTATTTCAGATTTCACCTTCCCTTATTTCAGTT

AV AV

E1A-F



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Amdt. dated February 20, 2004  
REPLACEMENT SHEET

Fig.10B.

| <- ITR

ATF

Wt : ::::::::::::::::::::::::::::::::::::::CTACGTCACCCGCCCGGTTCCACGCCCGCGG

Mut TTCCCGCGAAATGGCCAAATCTTACTCGGTTACGCCCAAATTTACTACTAGTGGAGATCAAAGGGATCCAGATCAAAGGGA

E2F AII

AI

Tcf

Tcf

| <- New ITR

ATF Spl Region of max DNA replication

Wt CCACGTCACAAACTCCACCCCTCATTATCATATTTGGCTTCAATCCCAAATAAGGTATATTATTGATGATG

Mut TTAAGATCAAAGGGCCACCACCTCATTATCATATTTGGCTTCAATCCCAAATAAGGTATATTATTGATGATG

Tcf



Fig.11.

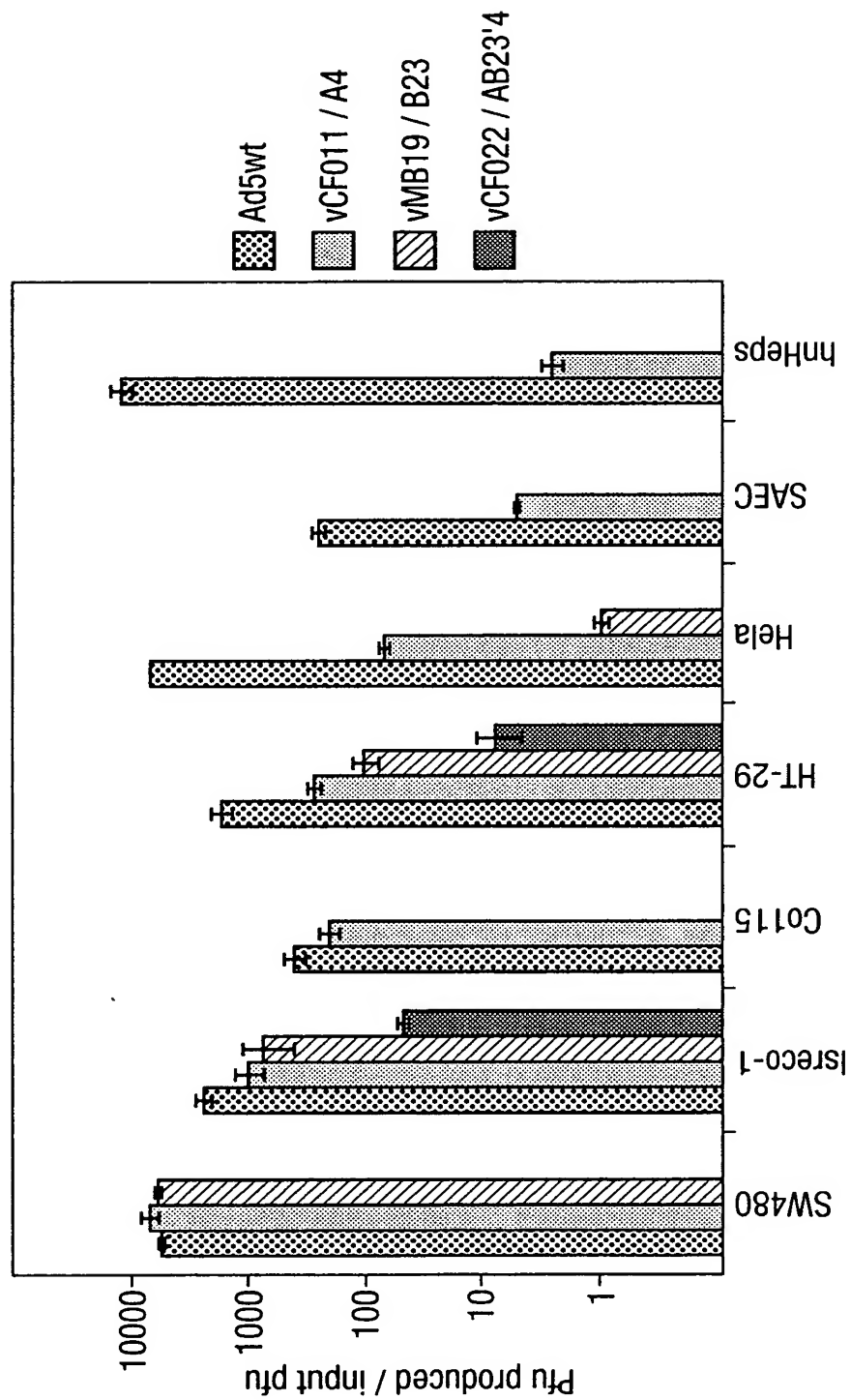
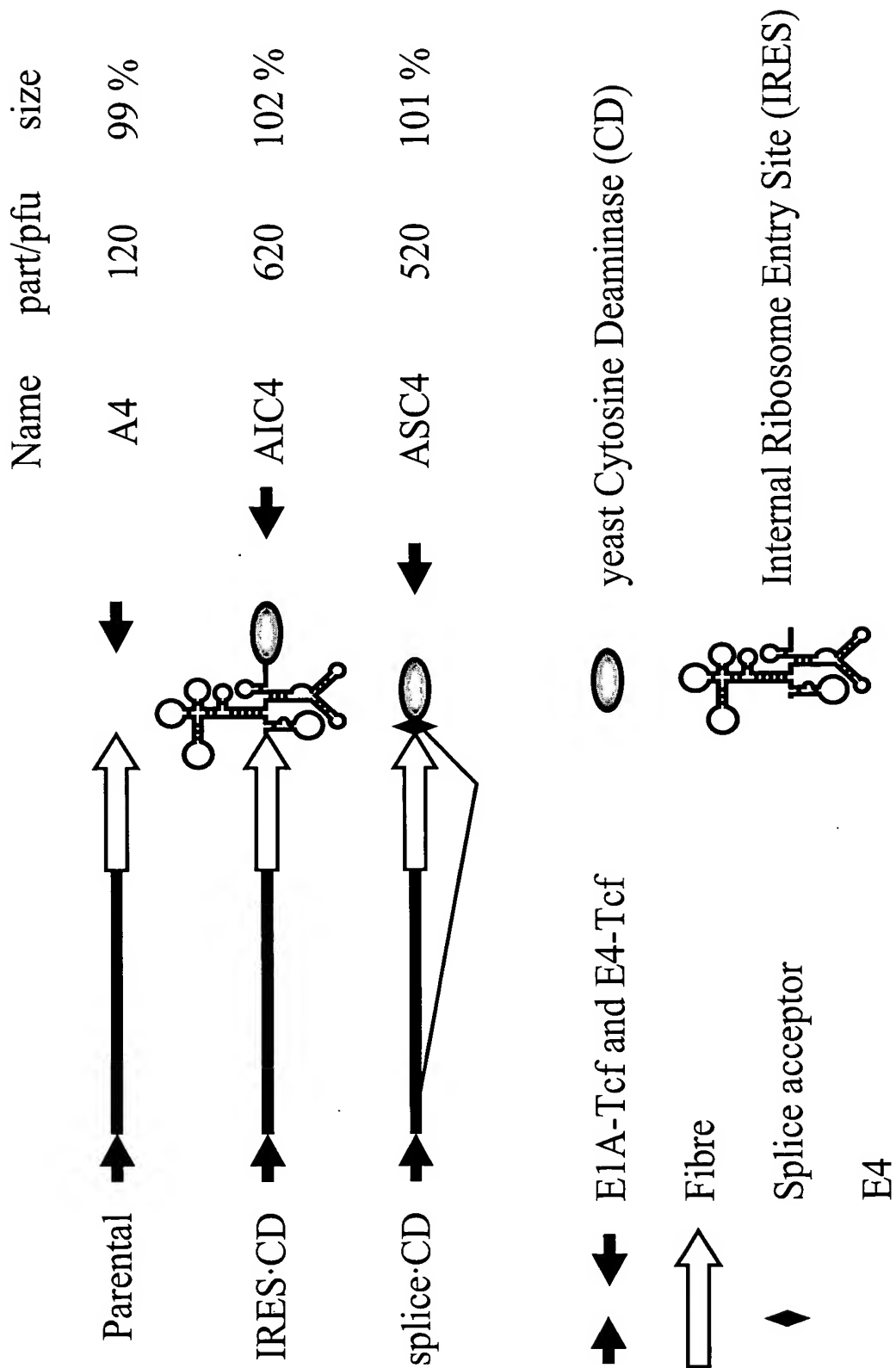


Fig.12.





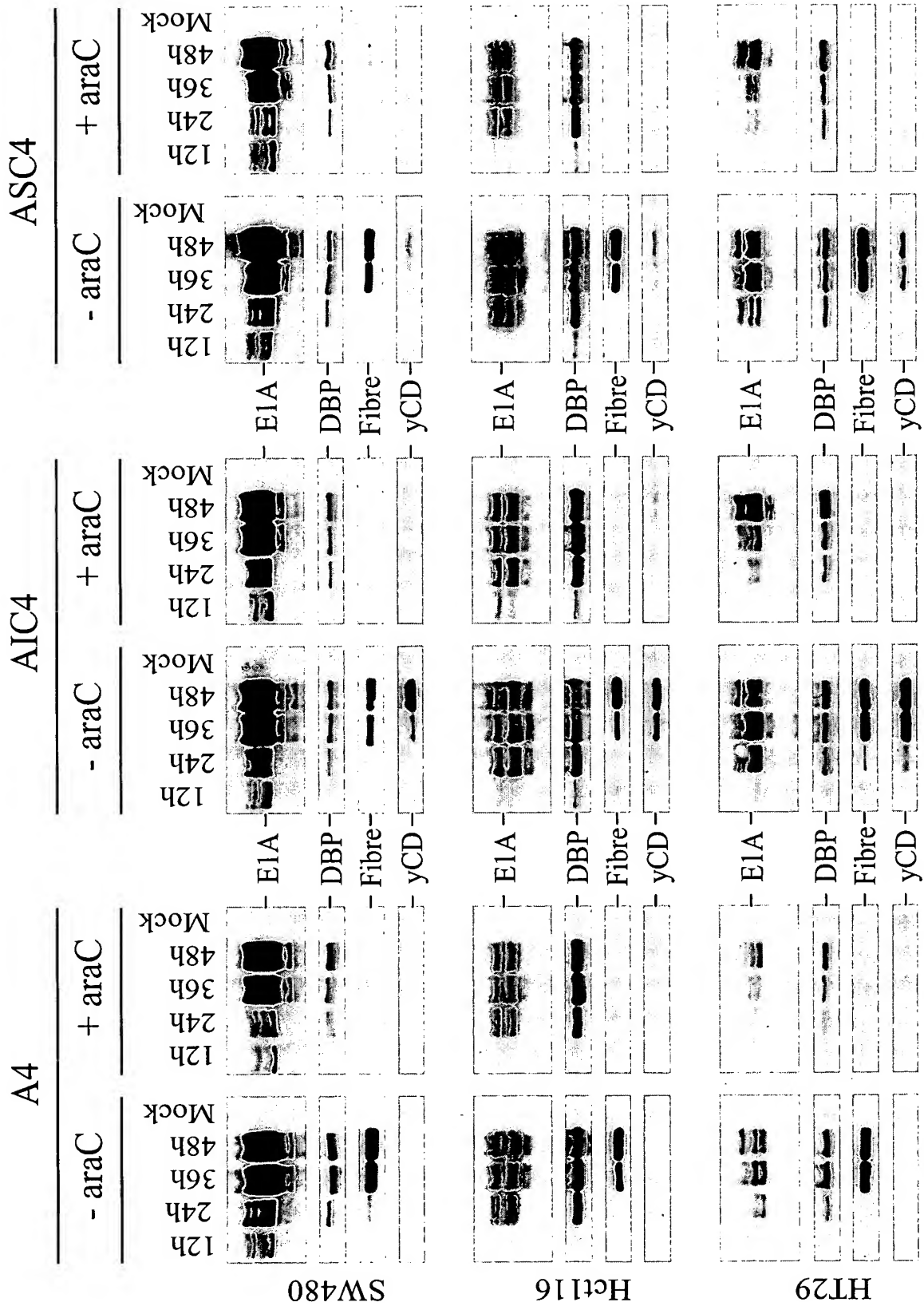


Fig.13.

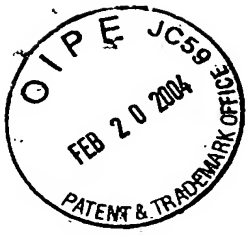


Fig. 14A.

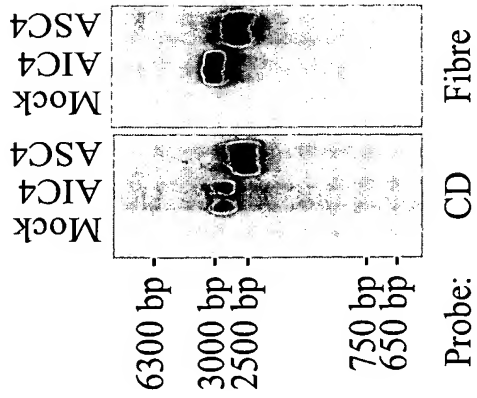


Fig. 14C.

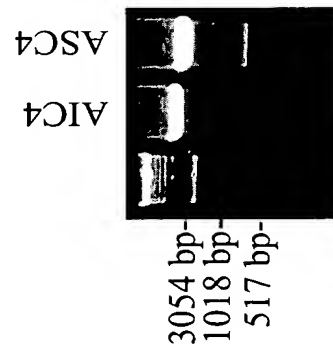
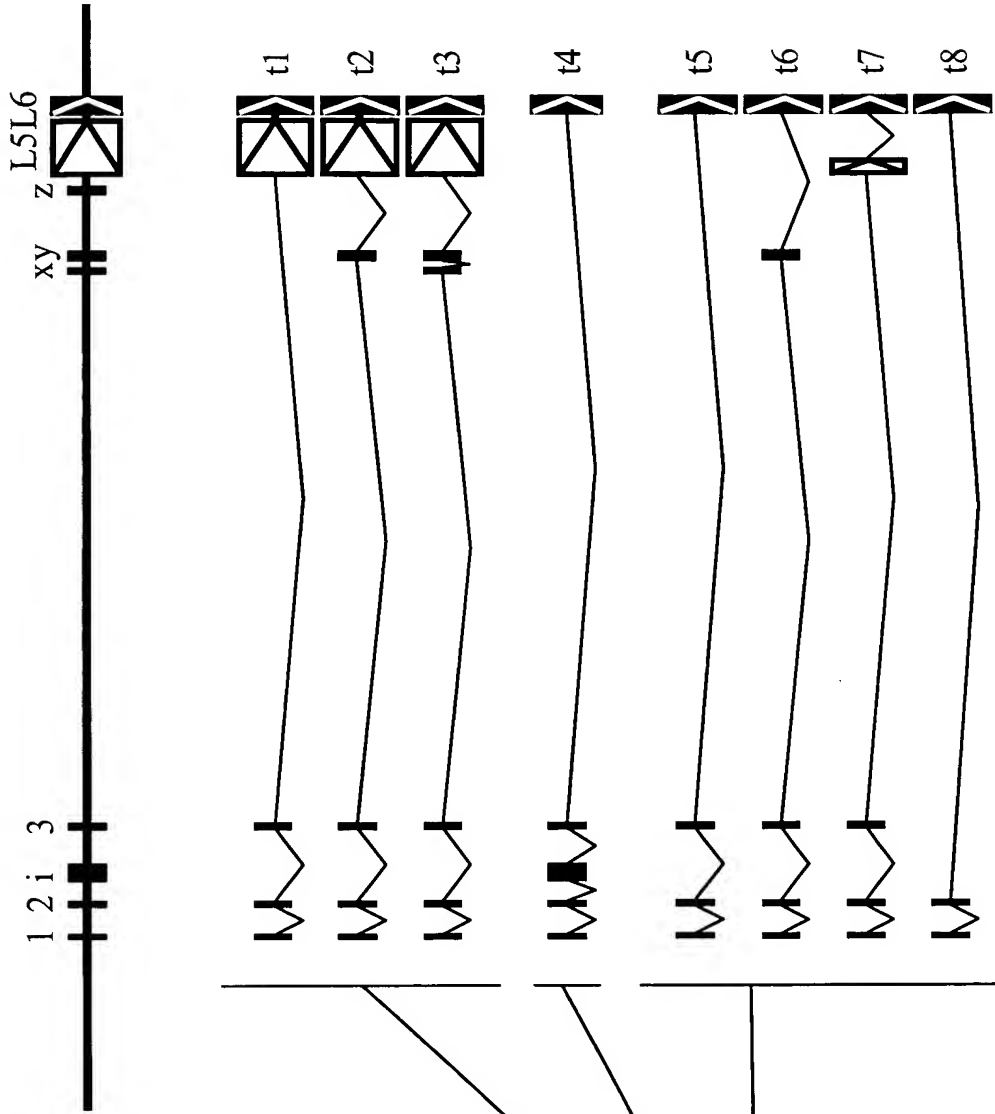


Fig. 14B.



Fig. 15A.

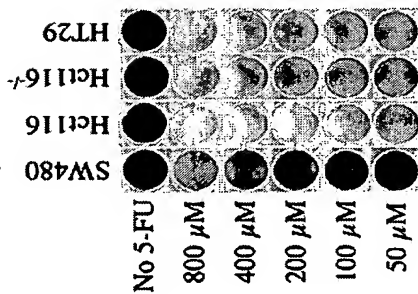


Fig. 15B.

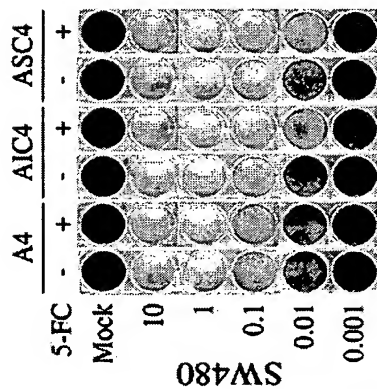


Fig. 15C.

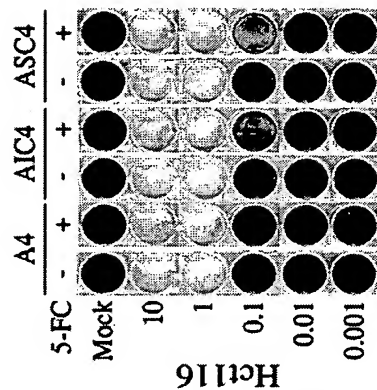


Fig. 15D.

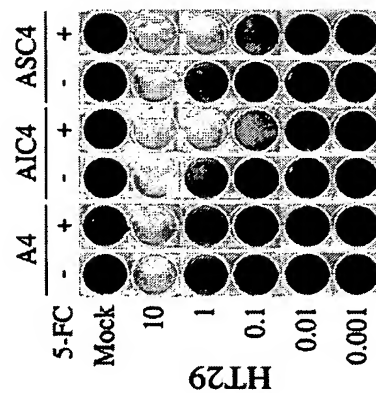
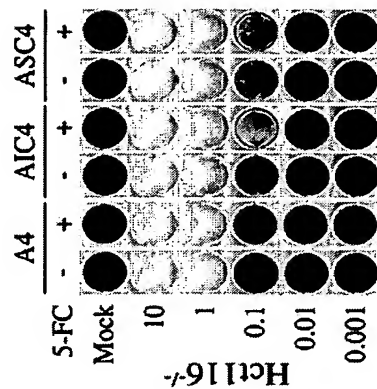


Fig. 15E.

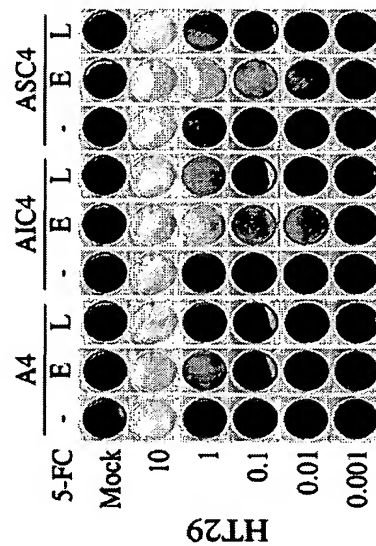


Fig. 15F.

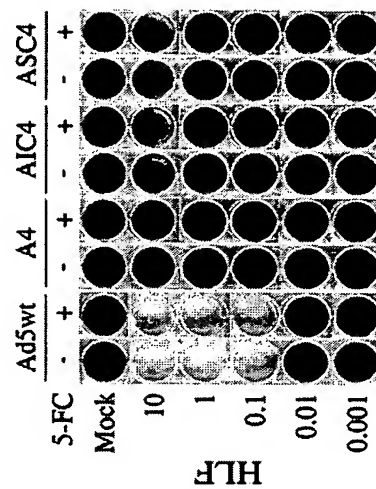


Fig. 15G.

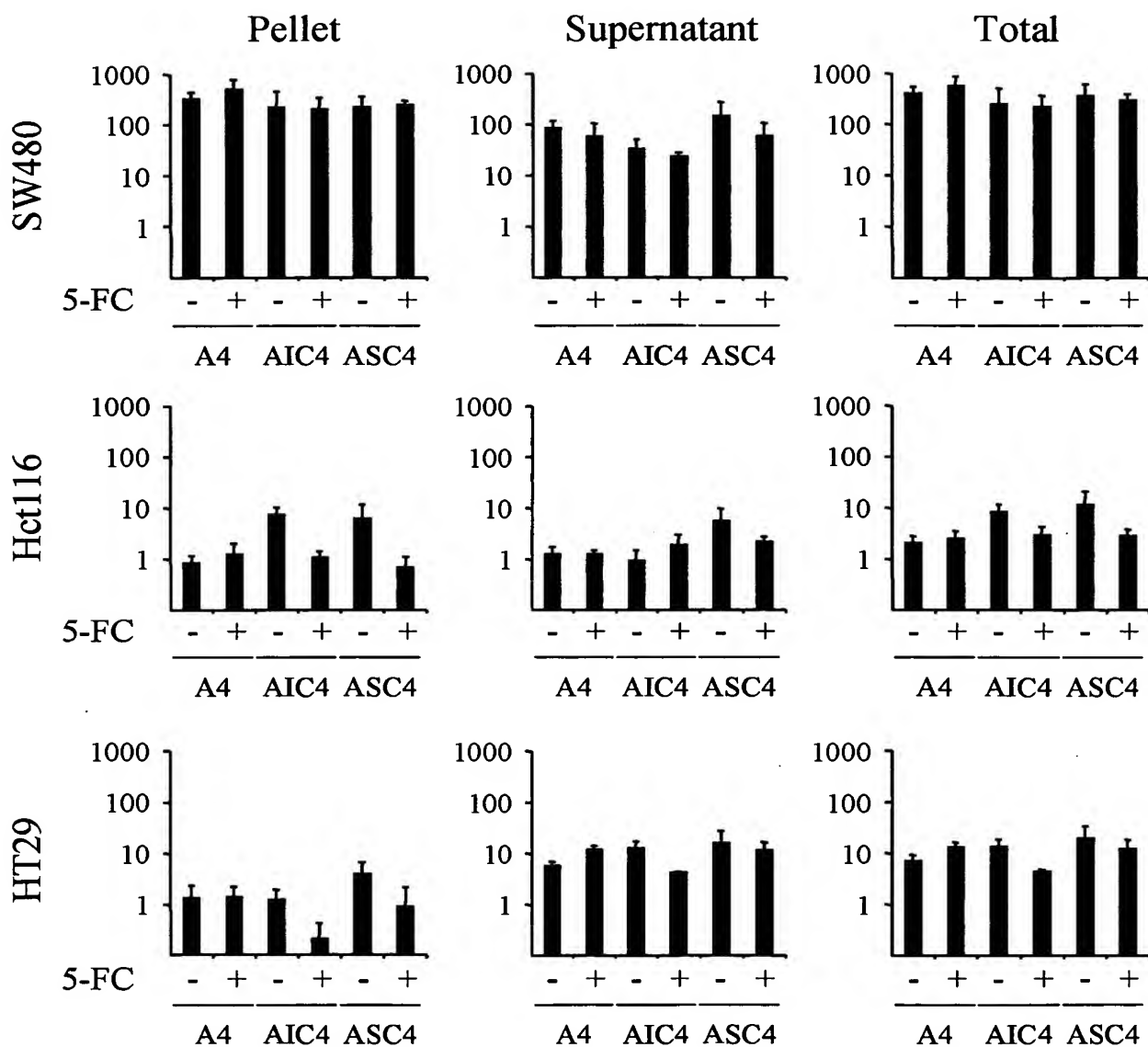


Fig.16.